CRF Errors Corrected by the STIC Systems Branch

Serial Number: _09/687,860_   **ENTERED**

CRF Processing Date: _10/27/2000_
Edited by: _____
Verified by: _____ (STIC staff)

- [ ] Changed a file from non-ASCII to ASCII

- [ ] Changed the margins in cases where the sequence text was "wrapped" down to the next line.

- [ ] Edited a format error in the Current Application Data section, specifically: _____

- [ ] Edited the Current Application Data section with the actual current number. The number inputted by the applicant was [ ] the prior application data; or [ ] other _____.

- [ ] Added the mandatory heading and subheadings for "Current Application Data".

- [ ] Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

- [ ] Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

- [ ] Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

- [x] Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _22_

- [ ] Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

- [ ] Inserted colons after headings/subheadings. Headings edited included: _____

- [ ] Deleted extra, invalid, headings used by an applicant, specifically: _____

- [ ] Deleted: [ ] non-ASCII "garbage" at the beginning/end of files; [ ] secretary initials/filename at end of file; [ ] page numbers throughout text; [ ] other invalid text, such as_____ .

- [ ] Inserted mandatory headings, specifically: _____

- [ ] Corrected an obvious error in the response, specifically: _____

- [ ] Edited identifiers where upper case is used but lower case is required, or vice versa.

- [ ] Corrected an error in the Number of Sequences field, specifically: _____

- [ ] A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

- [ ] Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

- [ ] Other: _____
_____
_____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

OIPE

RAW SEQUENCE LISTING   DATE: 10/27/2000
PATENT APPLICATION: US/09/687,860   TIME: 13:08:59

Input Set : A:\36737 sequence listing.txt
Output Set: N:\CRF3\10272000\I687860.raw

```
       3 <110> APPLICANT: Asundi, Vinod
       4        Ford, John E.
       5        Drmanac, Radoje T.
       6        Liu, Chenghua
       7        Tang, Y. Tom
       8        Yamasaki, Vicky
       9        Yeung, George
      10        Zhang, Jie
      11        Zhou, Ping
      13 <120> TITLE OF INVENTION: EGF MOTIF PROTEIN, EGFL6, MATERIALS AND METHODS
      15 <130> FILE REFERENCE: 28110/36737
C--> 17 <140> CURRENT APPLICATION NUMBER: US/09/687,860
C--> 17 <141> CURRENT FILING DATE: 2000-10-13
      17 <150> PRIOR APPLICATION NUMBER: US 09/620,312
      18 <151> PRIOR FILING DATE: 2000-07-19
      20 <150> PRIOR APPLICATION NUMBER: US 09/363,316
      21 <151> PRIOR FILING DATE: 1999-07-28
      23 <160> NUMBER OF SEQ ID NOS: 32
      25 <170> SOFTWARE: FastSEQ for Windows Version 3.0
```

ERRORED SEQUENCES

```
     841 <210> SEQ ID NO: 22
     842 <211> LENGTH: 20
     843 <212> TYPE: DNA
     844 <213> ORGANISM: Artificial Sequence
     846 <220> FEATURE:
     847 <223> OTHER INFORMATION: Description of Artificial Sequence: primer
     849 <400> SEQUENCE: 22
E--> 850 gggaactgac atacaaagtc                    20/-
```

VERIFICATION SUMMARY   DATE: 10/27/2000
PATENT APPLICATION: US/09/687,860   TIME: 13:09:01

Input Set : A:\36737 sequence listing.txt
Output Set: N:\CRF3\10272000\I687860.raw

L:17 M:270 C: Current Application Number differs, Replaced Current Application No
L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:61 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:62 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:103 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:104 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:207 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:308 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:416 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:5
L:416 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:5
L:535 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:574 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:576 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:592 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:594 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:596 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:639 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:641 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:643 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:661 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:805 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:850 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:22
L:1186 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1198 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1203 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1207 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1211 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1215 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1219 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1223 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1227 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1231 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1235 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1239 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1243 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1247 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1251 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1255 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1259 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1263 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1267 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1271 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1275 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1279 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1283 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1287 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27

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PATENT APPLICATION: US/09/687,860   TIME: 13:09:01

Input Set : A:\36737 sequence listing.txt
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```
L:1291 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1295 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1299 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1303 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1307 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1311 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1315 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1319 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1323 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1327 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1331 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1335 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27
L:1482 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:1494 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1498 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1502 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1506 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1510 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1514 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1518 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1522 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1526 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1530 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1534 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1538 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1542 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1546 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
L:1550 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29
```